

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/559,835
Source: JFWO
Date Processed by STIC: 12/20/2005

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. **EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)**
2. **U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450**
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314**

Revised 01/24/05



IFWO

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/559,835

DATE: 12/20/2005
TIME: 11:16:19

Input Set : A:\Sequence Listing filed 12-6-05.txt
Output Set: N:\CRF4\12202005\J559835.raw

1 <110> APPLICANT: KRINGLE PHARMA CO., LTD.
W--> 2 <110> APPLICANT: Nakamura, Toshikazu
W--> 3 <120> TITLE OF INVENTION: A pharmaceutical composition comprising cells
W--> 4 <130> FILE REFERENCE: K12F1262
C--> 5 <140> CURRENT APPLICATION NUMBER: US/10/559,835
C--> 5 <141> CURRENT FILING DATE: 2005-12-06
E--> 5 <160> NUMBER OF SEQ ID: 4 → found 6

Does Not Comply
Corrected Diskette Needed
(Pg -1)

ERRORED SEQUENCES

197 <210> SEQ ID NO: 5
198 <211> LENGTH: 20
E--> 199 <212> TYPE: → (Missing)
W--> 200 <213> ORGANISM: → 213? Responses are Artificial
W--> 201 <400> SEQUENCE: 5
202 agtaggtgg atggtagtt
204 <210> SEQ ID NO: 6
205 <211> LENGTH: 20
E--> 206 <212> TYPE: Artificial sequence
W--> 207 <213> ORGANISM: →
W--> 208 <400> SEQUENCE: 6
209 tacaacttgt atgtcaaaat

Responses are Artificial
Unknown or Genus species.
DIS See Item # 10 on
Error Summary Sheet.

→ Same Error 20
→ Same Error ,

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/559,835

DATE: 12/20/2005
TIME: 11:16:20

Input Set : A:\Sequence Listing filed 12-6-05.txt
Output Set: N:\CRF4\12202005\J559835.raw

L:2 M:280 W: Numeric Identifier already exists, <110> found multiple times
L:3 M:283 W: Missing Blank Line separator, <120> field identifier
L:4 M:283 W: Missing Blank Line separator, <130> field identifier
L:5 M:270 C: Current Application Number differs, Replaced Current Application No
L:5 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:5 M:283 W: Missing Blank Line separator, <160> field identifier
L:11 M:283 W: Missing Blank Line separator, <400> field identifier
L:44 M:283 W: Missing Blank Line separator, <400> field identifier
L:77 M:283 W: Missing Blank Line separator, <400> field identifier
L:139 M:283 W: Missing Blank Line separator, <400> field identifier
L:199 M:310 E: (3) Wrong or Missing Sequence Type, numeric identifier <212>, for
SEQ ID#:5
L:200 M:201 W: Mandatory field data missing, <213> ORGANISM
L:201 M:283 W: Missing Blank Line separator, <400> field identifier
L:206 M:310 E: (3) Wrong or Missing Sequence Type, numeric identifier <212>, for
SEQ ID#:6
L:207 M:201 W: Mandatory field data missing, <213> ORGANISM
L:208 M:283 W: Missing Blank Line separator, <400> field identifier
L:5 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (4) Counted (6)